

0570
1025 #2

OIPE

P.5

RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/960,481

DATE: 10/10/2001

TIME: 08:51:06

Input Set : D:\15367C seq listing.txt

Output Set: N:\CRF3\10102001\I960481.raw

1 <110> APPLICANT: Byrum, Joseph R.
 2 Heck, Gregory R.
 3 La Rosa, Thomas J.
 5 <120> TITLE OF INVENTION: Nucleic Acid Molecules And Other Molecules Associated With
 6 Plants
 8 <130> FILE REFERENCE: 38-21(15367)C
 C--> 10 <140> CURRENT APPLICATION NUMBER: US/09/960,481
 C--> 10 <141> CURRENT FILING DATE: 2001-09-24
 10 <150> PRIOR APPLICATION NUMBER: US 09/306,349
 11 <151> PRIOR FILING DATE: 1999-05-10
 13 <160> NUMBER OF SEQ ID NOS: 53893
 15 <210> SEQ ID NO: 1
 16 <211> LENGTH: 300
 17 <212> TYPE: DNA
 18 <213> ORGANISM: Glycine max
 20 <223> OTHER INFORMATION: Clone ID: 701036901H1
 22 <400> SEQUENCE: 1
 24 tgtaaatctc aagatcagat cctcatctac caccattcca ggtgtagcca tcaaaacaac 60
 26 gtttgtttaa caaaacccat tatcagatca gatccgatca gatcctcttc agattcggat 120
 28 caccgcccag tccatgagca cgcagcgcct ttcagggggc acgggcctct tctctggatt 180
 30 caccggttta tgcaagggcc ttgcggtggt gcttggttgc ggtcacattg cggttcagtt 240
 32 tatcccttca gctgttaact atcttgctct tattcccgcc aggacaatcc catttgctg 300
 35 <210> SEQ ID NO: 2
 36 <211> LENGTH: 297
 37 <212> TYPE: DNA
 38 <213> ORGANISM: Glycine max
 40 <223> OTHER INFORMATION: Clone ID: 701036902H1
 42 <400> SEQUENCE: 2
 44 aaattacata gctcaataa tctgagagag aaacaaaaac aatgaagggt tccattgcca 60
 46 agttccactc cctccccta tgttttttcc ttctcttggc cttcaacaca aaacccctcc 120
 48 tagcagcaga accagagcca gtggttgata agcaaggga ccctttggag ccaggagtag 180
 50 ggtactacgt gtggccactt tgggctgatg aaggaggcct cacactaggc caaacaagga 240
 52 acaagacatg cctcttttat gttatccgtg acccttcatt cattggcacc ccatttc 297
 55 <210> SEQ ID NO: 3
 56 <211> LENGTH: 295
 57 <212> TYPE: DNA
 58 <213> ORGANISM: Glycine max
 60 <220> FEATURE:
 61 <221> NAME/KEY: unsure
 62 <222> LOCATION: (1)..(295)
 63 <223> OTHER INFORMATION: unsure at all n locations
 64 <223> OTHER INFORMATION: Clone ID: 701036903H1
 66 <400> SEQUENCE: 3
 68 gcctagccat tgatttgga aagtccaagc ttcaaagatg agtaagcttc agagtgatgc 60
 W--> 70 tcttagagaa gctatctcan gaatcatgac tgattccaag gaaaagaaca ggaaatttgt 120
 72 ggagaccatt gaactccaaa ttgggctgaa aaactatgat cccagaagg acaagcgttt 180
 W--> 74 tantggtct gttaagttgc ctcacattcc tcgccccag atgaagattt gtatgcttgg 240

ENTERED

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76 agatgcccaa cacgttgaag aggctgagaa gataggattg gactacatgg atgtc      295
79 <210> SEQ ID NO: 4
80 <211> LENGTH: 291
81 <212> TYPE: DNA
82 <213> ORGANISM: Glycine max
84 <223> OTHER INFORMATION: Clone ID: 701036904H1
86 <400> SEQUENCE: 4
88 tttttcttct atgaacacat ggtggctgag ttgaggggaa gtacttagaa acagaaacac    60
90 agagatgaaa ccagaaacat ctctgaactt tactttgatg ctgactttct tactttatct    120
92 gaacttggtt atgggagtgct tgagcggtga tcattatcgc agagttgact ttcctgatga    180
94 attcgttttt gtttcaggca cctctgctta tcaggtggaa ggagcttcta acgaagatgg    240
96 gagaactcct agcatctggg atacctttgc ccatgcggtg tatgagcagg a              291
99 <210> SEQ ID NO: 5
100 <211> LENGTH: 292
101 <212> TYPE: DNA
102 <213> ORGANISM: Glycine max
104 <223> OTHER INFORMATION: Clone ID: 701036905H1
106 <400> SEQUENCE: 5
108 gtttttgccc accacttttg ctttttctgc tcaaattctg tgctcatcgc cctcgctcgtg    60
110 ttcactttgt cactgctaatt ggcatttttg ctggggaaac ggcgctcttc gcaaaacgaa    120
112 ttggcaaaac ggcgcagaga caaatcgcca gaaggcattg aattggagga gaatcccag    180
114 gtgttcaaaa gttctttgct gaaggactcc aaagaacaca tcatcagtac gtactttcct    240
116 gatgtgcgca cacttttcga tgctttccaa cgcggtctca aattgccgaa ca              292
119 <210> SEQ ID NO: 6
120 <211> LENGTH: 293
121 <212> TYPE: DNA
122 <213> ORGANISM: Glycine max
124 <220> FEATURE:
125 <221> NAME/KEY: unsure
126 <222> LOCATION: (1)..(293)
127 <223> OTHER INFORMATION: unsure at all n locations
128 <223> OTHER INFORMATION: Clone ID: 701036906H1
130 <400> SEQUENCE: 6
W--> 132 caaattttcg gatctcttct gtttcttcga atncaaaacc ctaattcccc caatttcatt    60
134 tttcaacctt gaaaaacctt ctctctctct ctgagttccc aacggtcac accacagctc    120
136 gatcaacagg gacatatattg ctactgaaga gatgtctcct gcttccaaat ctaagtctaa    180
138 agacaaaaat gctggcaagg aagctcaaaa gtcttctgcc aagtcttcag gatctggtaa    240
140 tgccgtggct ggtgtaccag ctagtgcata caatccttta tcgggaacat cca              293
143 <210> SEQ ID NO: 7
144 <211> LENGTH: 292
145 <212> TYPE: DNA
146 <213> ORGANISM: Glycine max
148 <223> OTHER INFORMATION: Clone ID: 701036907H1
150 <400> SEQUENCE: 7
152 ggcactcaaa cattgtttgt tgtaggaaga ggaatcttga gaaatggcag gggacagcaa    60
154 gagcaagatt ctgttcatcg gagggactgg ttacatcgga aaattcatcg tggaagccag    120
156 cgccaaagct ggccacccaa ctttcctttt ggtgagggag tccactctct ccaacccgc    180
158 aaaatcatcc ctcatcgaca atttcaaagg cctcggtgtc aatcttgttt tcggagatct    240
160 atacgatcac caaagtttgg tgagtgtctat aaagcaggtc gatgtggtaa tt              292

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163 <210> SEQ ID NO: 8
164 <211> LENGTH: 86
165 <212> TYPE: DNA
166 <213> ORGANISM: Glycine max
168 <220> FEATURE:
169 <221> NAME/KEY: unsure
170 <222> LOCATION: (1)..(86)
171 <223> OTHER INFORMATION: unsure at all n locations
172 <223> OTHER INFORMATION: Clone ID: 701036908H1
174 <400> SEQUENCE: 8
W--> 176 gaaaaanaaa aaaacngaac aactacngaa antcaaaacc aacttaaatt tnttatggat 60
W--> 178 ccannttnca tnggttangt tgtttt 86
181 <210> SEQ ID NO: 9
182 <211> LENGTH: 293
183 <212> TYPE: DNA
184 <213> ORGANISM: Glycine max
186 <220> FEATURE:
187 <221> NAME/KEY: unsure
188 <222> LOCATION: (1)..(293)
189 <223> OTHER INFORMATION: unsure at all n locations
190 <223> OTHER INFORMATION: Clone ID: 701036909H1
192 <400> SEQUENCE: 9
W--> 194 tcaaaccaaa aaacacaaca aaaganagan agaaaganac aaagagttcc ttgctttgta 60
W--> 196 gtaagatata tccatcaaaa atggttttgt cttggactag aaganacggt ttccgccgtg 120
W--> 198 cnaagaaggc gaaggaacaa agtcaactat tagaagttga ggttggtgatt cccaaccact 180
200 ttgcgtgtcc ggtgagtttg gagctgatga cagatccggt gactttatca accggcatta 240
202 cctacgaccg tgtgagcatc gagaagtgga tcgagggaga aaacagaacc tgc 293
205 <210> SEQ ID NO: 10
206 <211> LENGTH: 141
207 <212> TYPE: DNA
208 <213> ORGANISM: Glycine max
210 <220> FEATURE:
211 <221> NAME/KEY: unsure
212 <222> LOCATION: (1)..(141)
213 <223> OTHER INFORMATION: unsure at all n locations
214 <223> OTHER INFORMATION: Clone ID: 701036910H1
216 <400> SEQUENCE: 10
W--> 218 tggagtttta nttaaacta tggctatggt aatggaaaac cgncggttttt ncgtgttttc 60
W--> 220 accactttcg ttcttgatc agagtcaccc ccaaaaaaca aggcctaant cgccttcttc 120
W--> 222 ctctttccct tcttntttt t 141
225 <210> SEQ ID NO: 11
226 <211> LENGTH: 290
227 <212> TYPE: DNA
228 <213> ORGANISM: Glycine max
230 <220> FEATURE:
231 <221> NAME/KEY: unsure
232 <222> LOCATION: (1)..(290)
233 <223> OTHER INFORMATION: unsure at all n locations
234 <223> OTHER INFORMATION: Clone ID: 701036911H1

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Input Set : D:\15367C seq listing.txt

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236 <400> SEQUENCE: 11
238 ctcccgcatc tcctcgtcgc tccgccacgc cttcgcgag gccgaccggt ggacggagct 60
240 catcgaccgg agctccatgt cccgaccgga caccctcgcc gaagcatact ctcggtatccg 120
W--> 242 taagaacttc gcctanttcc gcgtcaatta cctaacccta atcgtgttgg cgctcgcgggt 180
244 ctcgctcatc acgcatccgt tctcgctatt cgtcctcttc ggtctcctcg cgctcgtggtc 240
W--> 246 gttcctttat ctctttcgcc cttcggatca gctgtcgtc ntcttcggac 290
249 <210> SEQ ID NO: 12
250 <211> LENGTH: 285
251 <212> TYPE: DNA
252 <213> ORGANISM: Glycine max
254 <220> FEATURE:
255 <221> NAME/KEY: unsure
256 <222> LOCATION: (1)..(285)
257 <223> OTHER INFORMATION: unsure at all n locations
258 <223> OTHER INFORMATION: Clone ID: 701036912H1
260 <400> SEQUENCE: 12
W--> 262 ggtgacgaga gccaccaccc accaccgcaa gcgcaantga aaaggttact gaagttggat 60
W--> 264 ggtacnttct tggcgaggat caacaacaga ttggccctta cgcttctct gagttgtgcc 120
266 agcatttctt gaatggatac ctctccgaaa atacgtttgt gtggtctgaa ggaagtagtg 180
268 agtggcagcc cttgtcctcg gtttctgacc tgtgggctca aattaaccgg caggacctga 240
270 ttcttcaact actgtgtctg ctctgatgt tgatgagttt gaaag 285
273 <210> SEQ ID NO: 13
274 <211> LENGTH: 289
275 <212> TYPE: DNA
276 <213> ORGANISM: Glycine max
278 <220> FEATURE:
279 <221> NAME/KEY: unsure
280 <222> LOCATION: (1)..(289)
281 <223> OTHER INFORMATION: unsure at all n locations
282 <223> OTHER INFORMATION: Clone ID: 701036913H1
284 <400> SEQUENCE: 13
286 cacttcttga accgtgcctt aactaaacta gagctagaat tagagttagc taccttgcct 60
W--> 288 aattcacaaa cgcgtccctc tacggctcta cctattagct atcttntttg tgctgtgatt 120
290 gaaattaatt tgtgatagct caccatggcc attgctactt gtgatcatga taagaacgca 180
W--> 292 aaggctttgc agttcattga ggatatgacc caaaacactg aaagtgtcca aganagggtc 240
W--> 294 cttgctgaga ttctcagcca aaactcccag accgaatacc tganacggt 289
297 <210> SEQ ID NO: 14
298 <211> LENGTH: 282
299 <212> TYPE: DNA
300 <213> ORGANISM: Glycine max
302 <220> FEATURE:
303 <221> NAME/KEY: unsure
304 <222> LOCATION: (1)..(282)
305 <223> OTHER INFORMATION: unsure at all n locations
306 <223> OTHER INFORMATION: Clone ID: 701036914H1
308 <400> SEQUENCE: 14
W--> 310 actgatcata aatcacatag ctagttgttt aaagtgagag anaaagaaag aagaanatgg 60
312 caaactttcc agttgttgac atggggaaac ttaacaccga agagagagca gcagccatgg 120
W--> 314 agattataaa anatgcctgt gagaactggg gtttctttga gttggtgaac catgggatat 180

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316 ccattgagtt gatggacact gtggagagggc tcacaaaaga gcactacaag aagactatgg 240
318 agcaaagggtt caaagaaatg gtggccagca aagggttgga gt 282
321 <210> SEQ ID NO: 15
322 <211> LENGTH: 290
323 <212> TYPE: DNA
324 <213> ORGANISM: Glycine max
326 <220> FEATURE:
327 <221> NAME/KEY: unsure
328 <222> LOCATION: (1)..(290)
329 <223> OTHER INFORMATION: unsure at all n locations
330 <223> OTHER INFORMATION: Clone ID: 701036915H1
332 <400> SEQUENCE: 15
334 ctaaagcaaa cccagttgtc aagggttgaa aagcttgata agatctaaga tttgaagttg 60
336 aagttccatg ccctgagatg aagttatgca caatttcacg atgaaatgat ccgagtttcc 120
W--> 338 ttcacatttt attggtgttc ttattatatt gttatatgtg gtgggtgact tgagtnattg 180
340 gacatctctg caatttgaag gactctttca tgtgattggt gaaaatatat atgttggcca 240
342 ctgttggttt taagtgcggt ttggtgaaag ttggttggtg ttgttagctt 290
345 <210> SEQ ID NO: 16
346 <211> LENGTH: 170
347 <212> TYPE: DNA
348 <213> ORGANISM: Glycine max
350 <223> OTHER INFORMATION: Clone ID: 701036916H1
352 <400> SEQUENCE: 16
354 gtttctctct acaagtccat atgtaaaatg gataacagaa agaaaaagaa aaatcacttt 60
356 ggtttgagaa ctttttaaag tttatattaa ctgtgttaag gttcataaaa ctgtagactg 120
358 atttgtgtga catgctccac agaaccttaa atttctctct attttgtcct 170
361 <210> SEQ ID NO: 17
362 <211> LENGTH: 302
363 <212> TYPE: DNA
364 <213> ORGANISM: Glycine max
366 <220> FEATURE:
367 <221> NAME/KEY: unsure
368 <222> LOCATION: (1)..(302)
369 <223> OTHER INFORMATION: unsure at all n locations
370 <223> OTHER INFORMATION: Clone ID: 701036917H1
372 <400> SEQUENCE: 17
W--> 374 ggccgtttcg aaagcnatga cgctatccac cgcgccctca ccgccaccgg cattagaaga 60
376 ggagccgtta cctcacgacg gcgacgatta cgacggcatc gtttcaactg acgacgaggg 120
W--> 378 tttcccgtn cgaattcccc gaatcaggaa cggcgctcga aaanagcagt ttcgccgagg 180
380 attcactatc gcagacggaa ccgattgggt cttcgctgtt gttgtcgctg gaagatagta 240
382 aacagagcga ttttagcgcaa gattcgggga agaagatgaa gcaggccaat ctgttccaga 300
384 tt 302
387 <210> SEQ ID NO: 18
388 <211> LENGTH: 294
389 <212> TYPE: DNA
390 <213> ORGANISM: Glycine max
392 <223> OTHER INFORMATION: Clone ID: 701036918H1
394 <400> SEQUENCE: 18
396 gtttccgata tccaagggga aatggcattg gtgttcaact accaactgga aataaaggaa 60

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→ Use of n and / or Xaa has been detected in the Sequence Listing. Review the Sequence Listing to ensure a corresponding explanation is present in the <220> to <223> fields of each sequence using n or Xaa.

VERIFICATION SUMMARY

PATENT APPLICATION: US/09/960,481

DATE: 10/10/2001

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Input Set : D:\15367C seq listing.txt

Output Set: N:\CRF3\10102001\I960481.raw

L:10 M:270 C: Current Application Number differs, Replaced Current Application No
L:10 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:70 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3
L:74 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3
L:132 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:6
L:176 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8
L:178 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:8
L:194 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:9
L:196 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:9
L:198 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:9
L:218 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:10
L:220 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:10
L:222 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:10
L:242 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:11
L:246 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:11
L:262 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:12
L:264 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:12
L:288 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:13
L:292 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:13
L:294 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:13
L:310 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:14
L:314 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:14
L:338 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:15
L:374 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:17
L:378 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:17
L:422 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:19
L:428 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:19
L:446 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:20
L:452 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:20
L:490 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:22
L:492 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:22
L:514 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:23
L:538 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:24
L:560 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:25
L:562 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:25
L:586 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:26
L:588 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:26
L:592 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:26
L:628 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:28
L:652 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:29
L:654 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:29
L:696 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:31
L:702 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:31
L:720 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:32
L:728 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:32
L:744 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:33
L:746 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:33
L:752 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:33

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L:772 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:34
L:792 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:35
L:816 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:36
L:820 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:36